

Sequence Listing

<110> Baker, Kevin
Botstein, David
Eaton, Dan
Ferrara, Napoleone
Filvaroff, Ellen
Gerritsen, Mary
Goddard, Audrey
Godowski, Paul
Grimaldi, Christopher
Gurney, Austin
Hillan, Kenneth
Kljavin, Ivar
Napier, Mary
Roy, Margaret
Tumas, Daniel
Wood, William

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Asp Asp Asp Asp Asp Glu Asp Asn Ser Leu Phe Pro Thr Arg Glu
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Pro Arg Ser His Phe Phe Pro Phe Asp Leu Phe Pro Met Cys Pro
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Gly Leu Thr Ser Val Pro Thr Asn Ile Pro Phe Asp Thr Arg Met
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Phe Phe Glu Gly Gln Gln Arg Pro His Gly Ala Arg Trp Ala Pro	710	715	720
Asn Tyr Asp Pro Leu Cys Ser Leu Cys Thr Cys Gln Arg Arg Thr	725	730	735
Val Ile Cys Asp Pro Val Val Cys Pro Pro Pro Ser Cys Pro His	740	745	750
Pro Val Gln Ala Pro Asp Gln Cys Cys Pro Val Cys Pro Glu Lys	755	760	765
Gln Asp Val Arg Asp Leu Pro Gly Leu Pro Arg Ser Arg Asp Pro	770	775	780
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Gly Thr Arg Trp His Pro Val Val Pro Pro Phe Gly Leu Ile Lys	800	805	810
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Glu Lys Val Gln Cys Pro Arg Leu Ala Cys Ala Gln Pro Val Arg	830	835	840
Val Asn Pro Thr Asp Cys Cys Lys Gln Cys Pro Val Gly Ser Gly	845	850	855
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 <213> Homo Sapien

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 Ser Ser Leu Ala Asn Pro Val Pro Ala Ala Pro Leu Ser Ala Pro
 35 40 45
 Gly Pro Cys Ala Ala Gln Pro Cys Arg Asn Gly Gly Val Cys Thr
 50 55 60
 Ser Arg Pro Glu Pro Asp Pro Gln His Pro Ala Pro Ala Gly Glu
 65 70 75
 Pro Gly Tyr Ser Cys Thr Cys Pro Ala Gly Ile Ser Gly Ala Asn
 80 85 90
 Cys Gln Leu Val Ala Asp Pro Cys Ala Ser Asn Pro Cys His His
 95 100 105
 Gly Asn Cys Ser Ser Ser Ser Ser Ser Ser Asp Gly Tyr Leu

110	115	120
Cys Ile Cys Asn Glu Gly Tyr Glu Gly	Pro Asn Cys Glu Gln Ala	
125	130	135
Leu Pro Ser Leu Pro Ala Thr Gly Trp	Thr Glu Ser Met Ala Pro	
140	145	150
Arg Gln Leu Gln Pro Val Pro Ala Thr	Gln Glu Pro Asp Lys Ile	
155	160	165
Leu Pro Arg Ser Gln Ala Thr Val Thr	Leu Pro Thr Trp Gln Pro	
170	175	180
Lys Thr Gly Gln Lys Val Val Glu Met	Lys Trp Asp Gln Val Glu	
185	190	195
Val Ile Pro Asp Ile Ala Cys Gly Asn	Ala Ser Ser Asn Ser Ser	
200	205	210
Ala Gly Gly Arg Leu Val Ser Phe Glu	Val Pro Gln Asn Thr Ser	
215	220	225
Val Lys Ile Arg Gln Asp Ala Thr Ala	Ser Leu Ile Leu Leu Trp	
230	235	240
Lys Val Thr Ala Thr Gly Phe Gln Gln	Cys Ser Leu Ile Asp Gly	
245	250	255
Arg Ser Val Thr Pro Leu Gln Ala Ser	Gly Gly Leu Val Leu Leu	
260	265	270
Glu Glu Met Leu Ala Leu Gly Asn Asn	His Phe Ile Gly Phe Val	
275	280	285
Asn Asp Ser Val Thr Lys Ser Ile Val	Ala Leu Arg Leu Thr Leu	
290	295	300
Val Val Lys Val Ser Thr Cys Val Pro	Gly Glu Ser His Ala Asn	
305	310	315
Asp Leu Glu Cys Ser Gly Lys Gly Lys	Cys Thr Thr Lys Pro Ser	
320	325	330
Glu Ala Thr Phe Ser Cys Thr Cys Glu	Glu Gln Tyr Val Gly Thr	
335	340	345
Phe Cys Glu Glu Tyr Asp Ala Cys Gln	Arg Lys Pro Cys Gln Asn	
350	355	360
Asn Ala Ser Cys Ile Asp Ala Asn Glu	Lys Gln Asp Gly Ser Asn	
365	370	375
Phe Thr Cys Val Cys Leu Pro Gly Tyr	Thr Gly Glu Leu Cys Gln	
380	385	390
Ser Lys Ile Asp Tyr Cys Ile Leu Asp	Pro Cys Arg Asn Gly Ala	
395	400	405

Demographic Data		Clinical Data		Outcome Data	
Variable	Value	Variable	Value	Variable	Value
Age (years)	65.2	Gender	Male	Survival (%)	78.5
Weight (kg)	72.5	Height (cm)	175.3	Quality of Life (VAS)	82.1
BMI (kg/m ²)	23.8	Heart Rate (b/min)	72.1	Healthcare Costs (€)	1250
Diabetes (%)	15.2	Blood Pressure (mmHg)	120/80	Medication (mg/day)	100
Cholesterol (%)	22.5	Smoking Status	Non-smoker	Compliance (%)	95.3
Alcohol Consumption (g/day)	20.1	Exercise Frequency (times/week)	3.2	Adverse Events (%)	5.1
Family History (%)	18.7	Previous Illnesses (%)	12.3	Long-term Outcomes (%)	85.4
Genetic Testing (%)	10.5	Current Medications (%)	8.9	Healthcare Utilization (%)	15.2
Insurance Type (%)	75.3	Healthcare Access (%)	92.1	Healthcare Satisfaction (%)	88.7
Education Level (%)	68.4	Healthcare Knowledge (%)	85.6	Healthcare Engagement (%)	79.2
Employment Status (%)	55.1	Healthcare Needs (%)	72.8	Healthcare Preferences (%)	65.4
Marital Status (%)	42.3	Healthcare Beliefs (%)	68.9	Healthcare Attitudes (%)	58.7
Religious Beliefs (%)	35.6	Healthcare Values (%)	62.1	Healthcare Ethics (%)	52.3
Cultural Background (%)	28.9	Healthcare Norms (%)	55.4	Healthcare Trends (%)	45.8
Socioeconomic Status (%)	22.1	Healthcare Policies (%)	48.7	Healthcare Research (%)	38.2
Healthcare System (%)	15.4	Healthcare Innovation (%)	42.3	Healthcare Future (%)	32.1
Healthcare Reform (%)	10.2	Healthcare Development (%)	35.6	Healthcare Progress (%)	25.4
Healthcare Policy (%)	8.7	Healthcare Legislation (%)	28.9	Healthcare Regulation (%)	18.7
Healthcare Regulation (%)	7.3	Healthcare Standards (%)	22.1	Healthcare Accreditation (%)	12.3
Healthcare Accreditation (%)	6.1	Healthcare Certification (%)	15.4	Healthcare Inspection (%)	8.9
Healthcare Inspection (%)	5.2	Healthcare Monitoring (%)	10.2	Healthcare Evaluation (%)	7.3
Healthcare Monitoring (%)	4.5	Healthcare Assessment (%)	8.7	Healthcare Review (%)	6.1
Healthcare Assessment (%)	3.8	Healthcare Audit (%)	7.3	Healthcare Feedback (%)	5.2
Healthcare Audit (%)	3.1	Healthcare Feedback (%)	6.1	Healthcare Improvement (%)	4.5
Healthcare Feedback (%)	2.5	Healthcare Improvement (%)	5.2	Healthcare Innovation (%)	3.8
Healthcare Improvement (%)	1.9	Healthcare Innovation (%)	4.5	Healthcare Research (%)	3.1
Healthcare Innovation (%)	1.3	Healthcare Research (%)	3.1	Healthcare Future (%)	2.5
Healthcare Research (%)	0.8	Healthcare Future (%)	2.5	Healthcare Progress (%)	1.9
Healthcare Future (%)	0.5	Healthcare Progress (%)	1.3	Healthcare Development (%)	0.8
Healthcare Development (%)	0.2	Healthcare Progress (%)	0.5	Healthcare Regulation (%)	0.2
Healthcare Progress (%)	0.1	Healthcare Development (%)	0.2	Healthcare Accreditation (%)	0.1
Healthcare Accreditation (%)	0.0	Healthcare Inspection (%)	0.1	Healthcare Monitoring (%)	0.0
Healthcare Inspection (%)	0.0	Healthcare Assessment (%)	0.0	Healthcare Audit (%)	0.0
Healthcare Assessment (%)	0.0	Healthcare Audit (%)	0.0	Healthcare Feedback (%)	0.0
Healthcare Audit (%)	0.0	Healthcare Feedback (%)	0.0	Healthcare Improvement (%)	0.0
Healthcare Feedback (%)	0.0	Healthcare Improvement (%)	0.0	Healthcare Innovation (%)	0.0
Healthcare Improvement (%)	0.0	Healthcare Innovation (%)	0.0	Healthcare Research (%)	0.0
Healthcare Innovation (%)	0.0	Healthcare Research (%)	0.0	Healthcare Future (%)	0.0
Healthcare Research (%)	0.0	Healthcare Future (%)	0.0	Healthcare Progress (%)	0.0
Healthcare Future (%)	0.0	Healthcare Progress (%)	0.0	Healthcare Development (%)	0.0
Healthcare Development (%)	0.0	Healthcare Progress (%)	0.0	Healthcare Regulation (%)	0.0
Healthcare Progress (%)	0.0	Healthcare Development (%)	0.0	Healthcare Accreditation (%)	0.0
Healthcare Accreditation (%)	0.0	Healthcare Inspection (%)	0.0	Healthcare Monitoring (%)	0.0
Healthcare Inspection (%)	0.0	Healthcare Assessment (%)	0.0	Healthcare Audit (%)	0.0
Healthcare Assessment (%)	0.0	Healthcare Audit (%)	0.0	Healthcare Feedback (%)	0.0
Healthcare Audit (%)	0.0	Healthcare Feedback (%)	0.0	Healthcare Improvement (%)	0.0
Healthcare Feedback (%)	0.0	Healthcare Improvement (%)	0.0	Healthcare Innovation (%)	0.0
Healthcare Improvement (%)	0.0	Healthcare Innovation (%)	0.0	Healthcare Research (%)	0.0
Healthcare Innovation (%)	0.0	Healthcare Research (%)	0.0	Healthcare Future (%)	0.0
Healthcare Research (%)	0.0	Healthcare Future (%)	0.0	Healthcare Progress (%)	0.0
Healthcare Future (%)	0.0	Healthcare Progress (%)	0.0	Healthcare Development (%)	0.0
Healthcare Development (%)	0.0	Healthcare Progress (%)	0.0	Healthcare Regulation (%)	0.0
Healthcare Progress (%)	0.0	Healthcare Development (%)	0.0	Healthcare Accreditation (%)	0.0
Healthcare Accreditation (%)	0.0	Healthcare Inspection (%)	0.0	Healthcare Monitoring (%)	0.0
Healthcare Inspection (%)	0.0	Healthcare Assessment (%)	0.0	Healthcare Audit (%)	0.0
Healthcare Assessment (%)	0.0	Healthcare Audit (%)	0.0	Healthcare Feedback (%)	0.0
Healthcare Audit (%)	0.0	Healthcare Feedback (%)	0.0	Healthcare Improvement (%)	0.0
Healthcare Feedback (%)	0.0	Healthcare Improvement (%)	0.0	Healthcare Innovation (%)	0.0
Healthcare Improvement (%)	0.0	Healthcare Innovation (%)	0.0	Healthcare Research (%)	0.0
Healthcare Innovation (%)	0.0	Healthcare Research (%)	0.0	Healthcare Future (%)	0.0
Healthcare Research (%)	0.0	Healthcare Future (%)	0.0	Healthcare Progress (%)	0.0
Healthcare Future (%)	0.				

695

700

705

Ser Arg Pro Ala Met Tyr Asp Val Ser Pro Ile Ala Tyr Glu Asp
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Tyr Ser Pro Asp Asp Lys Pro Leu Val Thr Leu Ile Lys Thr Lys
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Asp Leu

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<213> Artificial Sequence

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<223> Synthetic Oligonucleotide Probe

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<213> Artificial Sequence

<220>

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<210> 18

<211> 508

<212> DNA

<213> Homo Sapien

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tctgtgacta agtctattgt ggctttgcgc ttaactctgg tggatgaagg 200

cagcacctgt gtgcggggg agagtcacgc aaatgacttg gagggttcag 250

gaaaaggaaa atgcaccacg aagccgtcag aggcaacttt ttctgtacc 300

tgtgaggagc agtacgtggg tactttctgt gaagaatacg atgcttgcca 350

gaggaaacct tgccaaaaca acgcgagctg tattgatgca aatgaaaagc 400

aagatgggag caatttcacc tgtgtttgcc ttctgggtta tactggagag 450

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taggggag 508

<210> 19

<211> 508

<212> DNA

<213> Homo Sapien

<400> 19

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tctgtgacta agtctattgt ggctttgcgc ttaactctgg tggatgaagg 200

cagcacctgt gtgcggggg agagtcacgc aaatgacttg gagggttcag 250

gaaaaggaaa atgcaccacg aagccgtcag aggcaacttt ttctgtacc 300

tgtgaggagc agtacgtggg tactttctgt gaagaatacg atgcttgcca 350

gaggaaacct tgccaaaaca acgcgagctg tattgatgca aatgaaaagc 400

aagatgggag caatttcacc tgtgtttgcc ttctgggta tactggagag 450

ctttgccaac cgaactgaga ttggagcgaa cgacctacac cgaactgaga 500

taggggag 508

<210> 20

<211> 23

<212> DNA

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<220>

<223> Synthetic Oligonucleotide Probe

<400> 20

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<210> 21

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

<400> 21

ctcagttcgg ttggcaaagc tctc 24

<210> 22

<211> 69

<212> DNA

<213> Artificial Sequence

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<211> 1520

<212> DNA

<213> Homo Sapien

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cagcaccag ggcctgcaag agcagggcacg ggcctgatg cgggacttcc 200

cgctcgtgga cggccacaac gacctgcccc tggtcctaag qcaagtttac 250

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cagcctggac aggccttagag atggcctcgt gggcgcccag ttctggtcag 350

cctatgtgcc atgccagacc caggaccggg atgccctgcg cctcaccctg 400

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<211> 433

<213> Homo Sapien

<213> Homo Sapien

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35 40 45

Val Tyr Gln Lys Gly Leu Gln Asp Val Asn Leu Arg Asn Phe Ser
50 55 60

Tyr Gly Gln Thr Ser Leu Asp Arg Leu Arg Asp Gly Leu Val Gly
65 70 75

Ala Gln Phe Trp Ser Ala Tyr Val Pro Cys Gln Thr Gln Asp Arg
80 85 90

Asp Ala Leu Arg Leu Thr Leu Glu Gln Ile Asp Leu Ile Arg Arg
95 100 105

Met Cys Ala Ser Tyr Ser Glu Leu Glu Leu Val Thr Ser Ala Lys
110 115 120

Ala Leu Asn Asp Thr Gln Lys Leu Ala Cys Leu Ile Gly Val Glu
125 130 135

Gly Gly His Ser Leu Asp Asn Ser Leu Ser Ile Leu Arg Thr Phe
140 145 150

Tyr Met Leu Gly Val Arg Tyr Leu Thr Leu Thr His Thr Cys Asn
155 160 165

Thr Pro Trp Ala Glu Ser Ser Ala Lys Gly Val His Ser Phe Tyr
170 175 180

Asn	Asn	Ile	Ser	Gly 185	Leu	Thr	Asp	Phe	Gly 190	Glu	Lys	Val	Val	Ala 195
Glu	Met	Asn	Arg	Leu 200	Gly	Met	Met	Val	Asp 205	Leu	Ser	His	Val	Ser 210
Asp	Ala	Val	Ala	Arg 215	Arg	Ala	Leu	Glu	Val 220	Ser	Gln	Ala	Pro	Val 225
Ile	Phe	Ser	His	Ser 230	Ala	Ala	Arg	Gly	Val 235	Cys	Asn	Ser	Ala	Arg 240
Asn	Val	Pro	Asp	Asp 245	Ile	Leu	Gln	Leu	Leu 250	Lys	Lys	Asn	Gly	Gly 255
Val	Val	Met	Val	Ser 260	Leu	Ser	Met	Gly	Val 265	Ile	Gln	Cys	Asn	Pro 270
Ser	Ala	Asn	Val	Ser 275	Thr	Val	Ala	Asp	His 280	Phe	Asp	His	Ile	Lys 285
Ala	Val	Ile	Gly	Ser 290	Lys	Phe	Ile	Gly	Ile 295	Gly	Gly	Asp	Tyr	Asp 300
Gly	Ala	Gly	Lys	Phe 305	Pro	Gln	Gly	Leu	Glu 310	Asp	Val	Ser	Thr	Tyr 315
Pro	Val	Leu	Ile	Glu 320	Glu	Leu	Leu	Ser	Arg 325	Gly	Trp	Ser	Glu	Glu 330
Glu	Leu	Gln	Gly	Val 335	Leu	Arg	Gly	Asn	Leu 340	Leu	Arg	Val	Phe	Arg 345
Gln	Val	Glu	Lys	Val 350	Gln	Glu	Glu	Asn	Lys 355	Trp	Gln	Ser	Pro	Leu 360
Glu	Asp	Lys	Phe	Pro 365	Asp	Glu	Gln	Leu	Ser 370	Ser	Ser	Cys	His	Ser 375
Asp	Leu	Ser	Arg	Leu 380	Arg	Gln	Arg	Gln	Ser 385	Leu	Thr	Ser	Gly	Gln 390
Glu	Leu	Thr	Glu	Ile 395	Pro	Ile	His	Trp	Thr 400	Ala	Lys	Leu	Pro	Ala 405
Lys	Trp	Ser	Val	Ser 410	Glu	Ser	Ser	Pro	His 415	Met	Ala	Pro	Val	Leu 420
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<212> DNA

<213> Artificial Sequence

<223> Synthetic oligonucleotide probe

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<210> 26
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<210> 27
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<212> DNA
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<223> Synthetic oligonucleotide probe

<400> 27
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<210> 28
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<212> DNA
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<220>
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<212> DNA
<213> Homo Sapien

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gccctgatgc gggacttccc gtcgtggac ggccacaacg acctgcccct 200
ggtcctaagg cagggttacc agaaagggt acaggatgtt aacctgcgca 250
atttcagcta cggccagacc agcctggaca ggcttagaga tggcctcgtg 300
ggcgcccagt tctggtcagc ctatgtgccca tgccagaccc aggaccggga 350
tgccctgcgc ctcaccctgg agcagattga cctcatagc cgcgatgtgtg 400

	50		55		60
Tyr Gly Gln Thr	Ser Leu Asp Arg Leu Arg Asp Gly Leu Val Gly				
	65		70		75
Ala Gln Phe Trp	Ser Ala Tyr Val Pro Cys Gln Thr Gln Asp Arg				
	80		85		90
Asp Ala Leu Arg	Leu Thr Leu Glu Gln Ile Asp Leu Ile Arg Arg				
	95		100		105
Met Cys Ala Ser	Tyr Ser Glu Leu Glu Leu Val Thr Ser Ala Lys				
	110		115		120
Ala Leu Asn Asp	Thr Gln Lys Leu Ala Cys Leu Ile Gly Val Glu				
	125		130		135
Gly Gly His Ser	Leu Asp Asn Ser Leu Ser Ile Leu Arg Thr Phe				
	140		145		150
Tyr Met Leu Gly	Val Arg Tyr Leu Thr Leu Thr His Thr Cys Asn				
	155		160		165
Thr Pro Trp Ala	Glu Ser Ser Ala Lys Gly Val His Ser Phe Tyr				
	170		175		180
Asn Asn Ile Ser	Gly Leu Thr Asp Phe Gly Glu Lys Val Val Ala				
	185		190		195
Glu Met Asn Arg	Leu Gly Met Met Val Asp Leu Ser His Val Ser				
	200		205		210
Asp Ala Val Ala	Arg Arg Ala Leu Glu Val Ser Gln Ala Pro Val				
	215		220		225
Ile Phe Ser His	Ser Ala Ala Arg Gly Val Cys Asn Ser Ala Arg				
	230		235		240
Asn Val Pro Asp	Asp Ile Leu Gln Leu Leu Lys Lys Asn Gly Gly				
	245		250		255
Val Val Met Val	Ser Leu Ser Met Gly Val Ile Gln Cys Asn Pro				
	260		265		270
Ser Ala Asn Val	Ser Thr Val Ala Asp His Phe Asp His Ile Lys				
	275		280		285
Ala Val Ile Gly	Ser Lys Phe Ile Gly Ile Gly Gly Asp Tyr Asp				
	290		295		300
Gly Ala Gly Lys	Phe Pro Gln Gly Leu Glu Asp Val Ser Thr Tyr				
	305		310		315
Pro Val Leu Ile	Glu Glu Leu Leu Ser Arg Gly Trp Ser Glu Glu				
	320		325		330
Glu Leu Gln Gly	Val Leu Arg Gly Asn Leu Leu Arg Val Phe Arg				
	335		340		345

Variable	Mean	SD	Min	Max
Age	34.5	10.5	20	55
Gender	1.0	0.0	0	1
Marital status	1.0	0.0	0	1
Education	12.5	1.5	9	16
Income	1.0	0.0	0	1
Health status	1.0	0.0	0	1
Smoking status	1.0	0.0	0	1
Alcohol consumption	1.0	0.0	0	1
Exercise frequency	1.0	0.0	0	1
Stress level	1.0	0.0	0	1
Sleep quality	1.0	0.0	0	1
Appetite	1.0	0.0	0	1
Energy level	1.0	0.0	0	1
Mood	1.0	0.0	0	1
Concentration	1.0	0.0	0	1
Memory	1.0	0.0	0	1
Emotional stability	1.0	0.0	0	1
Resilience	1.0	0.0	0	1
Optimism	1.0	0.0	0	1
Gratitude	1.0	0.0	0	1
Forgiveness	1.0	0.0	0	1
Compassion	1.0	0.0	0	1
Patience	1.0	0.0	0	1
Kindness	1.0	0.0	0	1
Generosity	1.0	0.0	0	1
Humility	1.0	0.0	0	1
Modesty	1.0	0.0	0	1
Shyness	1.0	0.0	0	1
Introversion	1.0	0.0	0	1
Extroversion	1.0	0.0	0	1
Sensitiveness	1.0	0.0	0	1
Emotional sensitivity	1.0	0.0	0	1
Empathy	1.0	0.0	0	1
Empathetic concern	1.0	0.0	0	1
Compassionate love	1.0	0.0	0	1
Self-love	1.0	0.0	0	1
Love for others	1.0	0.0	0	1
Love for life	1.0	0.0	0	1
Love for nature	1.0	0.0	0	1
Love for art	1.0	0.0	0	1
Love for music	1.0	0.0	0	1
Love for food	1.0	0.0	0	1
Love for travel	1.0	0.0	0	1
Love for learning	1.0	0.0	0	1
Love for growth	1.0	0.0	0	1
Love for change	1.0	0.0	0	1
Love for adventure	1.0	0.0	0	1
Love for risk	1.0	0.0	0	1
Love for challenge	1.0	0.0	0	1
Love for achievement	1.0	0.0	0	1
Love for success	1.0	0.0	0	1
Love for power	1.0	0.0	0	1
Love for status	1.0	0.0	0	1
Love for wealth	1.0	0.0	0	1
Love for fame	1.0	0.0	0	1
Love for recognition	1.0	0.0	0	1
Love for respect	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0.0	0	1
Love for admiration	1.0	0.0	0	1
Love for praise	1.0	0.0	0	1
Love for compliments	1.0	0.0	0	1
Love for attention	1.0	0		

<211> 1790

<212> DNA

<213> Homo Sapien

<400> 31

27

Ser Val His Gly Asp	Pro Pro Gly Ala Thr Ala Glu Gly Leu Tyr	65	70	75
Trp Thr Leu Asn Gly Arg Arg Leu Pro	Pro Glu Leu Ser Arg Val	80	85	90
Leu Asn Ala Ser Thr Leu Ala Leu Ala	Leu Ala Asn Leu Asn Gly	95	100	105
Ser Arg Gln Arg Ser Gly Asp Asn Leu	Val Cys His Ala Arg Asp	110	115	120
Gly Ser Ile Leu Ala Gly Ser Cys Leu	Tyr Val Gly Leu Pro Pro	125	130	135
Glu Lys Pro Val Asn Ile Ser Cys Trp	Ser Lys Asn Met Lys Asp	140	145	150
Leu Thr Cys Arg Trp Thr Pro Gly Ala	His Gly Glu Thr Phe Leu	155	160	165
His Thr Asn Tyr Ser Leu Lys Tyr Lys	Leu Arg Trp Tyr Gly Gln	170	175	180
Asp Asn Thr Cys Glu Glu Tyr His Thr	Val Gly Pro His Ser Cys	185	190	195
His Ile Pro Lys Asp Leu Ala Leu Phe	Thr Pro Tyr Glu Ile Trp	200	205	210
Val Glu Ala Thr Asn Arg Leu Gly Ser	Ala Arg Ser Asp Val Leu	215	220	225
Thr Leu Asp Ile Leu Asp Val Val Thr	Thr Asp Pro Pro Pro Asp	230	235	240
Val His Val Ser Arg Val Gly Gly Leu	Glu Asp Gln Leu Ser Val	245	250	255
Arg Trp Val Ser Pro Pro Ala Leu Lys	Asp Phe Leu Phe Gln Ala	260	265	270
Lys Tyr Gln Ile Arg Tyr Arg Val Glu	Asp Ser Val Asp Trp Lys	275	280	285
Val Val Asp Asp Val Ser Asn Gln Thr	Ser Cys Arg Leu Ala Gly	290	295	300
Leu Lys Pro Gly Thr Val Tyr Phe Val	Gln Val Arg Cys Asn Pro	305	310	315
Phe Gly Ile Tyr Gly Ser Lys Lys Ala	Gly Ile Trp Ser Glu Trp	320	325	330
Ser His Pro Thr Ala Ala Ser Thr Pro	Arg Ser Glu Arg Pro Gly	335	340	345
Pro Gly Gly Gly Ala Cys Glu Pro Arg	Gly Gly Glu Pro Ser Ser			

350	355	360
Gly Pro Val Arg Arg Glu Leu Lys Gln Phe Leu Gly Trp Leu Lys		
365	370	375
Lys His Ala Tyr Cys Ser Asn Leu Ser Phe Arg Leu Tyr Asp Gln		
380	385	390
Trp Arg Ala Trp Met Gln Lys Ser His Lys Thr Arg Asn Gln Asp		
395	400	405
Glu Gly Ile Leu Pro Ser Gly Arg Arg Gly Thr Ala Arg Gly Pro		
410	415	420
Ala Arg		

<210> 33
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 33
 cccgcccgcac gtgcacgtga gcc 23

<210> 34
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 34
 tgagccagcc caggaactgc ttg 23

<210> 35
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 35
 caagtgcgct gcaacccctt tggcatctat ggctccaaga aagccgggat 50

<210> 36
 <211> 1771
 <212> DNA
 <213> Homo Sapien

<400> 36
 cccacgcgtc cgctggtgtt agatcgagca accctctaaa agcagtttag 50

Sociodemographic variables		Sociodemographic variables	
Age	Gender	Age	Gender
18-24	Male	25-34	Male
25-34	Female	35-44	Female
35-44	Male	45-54	Male
45-54	Female	55-64	Female
55-64	Male	65-74	Male
65-74	Female	75-84	Female
75-84	Male	85-94	Male
85-94	Female	95-104	Female
95-104	Male	105-114	Male
105-114	Female	115-124	Female
115-124	Male	125-134	Male
125-134	Female	135-144	Female
135-144	Male	145-154	Male
145-154	Female	155-164	Female
155-164	Male	165-174	Male
165-174	Female	175-184	Female
175-184	Male	185-194	Male
185-194	Female	195-204	Female
195-204	Male	205-214	Male
205-214	Female	215-224	Female
215-224	Male	225-234	Male
225-234	Female	235-244	Female
235-244	Male	245-254	Male
245-254	Female	255-264	Female
255-264	Male	265-274	Male
265-274	Female	275-284	Female
275-284	Male	285-294	Male
285-294	Female	295-304	Female
295-304	Male	305-314	Male
305-314	Female	315-324	Female
315-324	Male	325-334	Male
325-334	Female	335-344	Female
335-344	Male	345-354	Male
345-354	Female	355-364	Female
355-364	Male	365-374	Male
365-374	Female	375-384	Female
375-384	Male	385-394	Male
385-394	Female	395-404	Female
395-404	Male	405-414	Male
405-414	Female	415-424	Female
415-424	Male	425-434	Male
425-434	Female	435-444	Female
435-444	Male	445-454	Male
445-454	Female	455-464	Female
455-464	Male	465-474	Male
465-474	Female	475-484	Female
475-484	Male	485-494	Male
485-494	Female	495-504	Female
495-504	Male	505-514	Male
505-514	Female	515-524	Female
515-524	Male	525-534	Male
525-534	Female	535-544	Female
535-544	Male	545-554	Male
545-554	Female	555-564	Female
555-564	Male	565-574	Male
565-574	Female	575-584	Female
575-584	Male	585-594	Male
585-594	Female	595-604	Female
595-604	Male	605-614	Male
605-614	Female	615-624	Female
615-624	Male	625-634	Male
625-634	Female	635-644	Female
635-644	Male	645-654	Male
645-654	Female	655-664	Female
655-664	Male	665-674	Male
665-674	Female	675-684	Female
675-684	Male	685-694	Male
685-694	Female	695-704	Female
695-704	Male	705-714	Male
705-714	Female	715-724	Female
715-724	Male	725-734	Male
725-734	Female	735-744	Female
735-744	Male	745-754	Male
745-754	Female	755-764	Female
755-764	Male	765-774	Male
765-774	Female	775-784	Female
775-784	Male	785-794	Male
785-794	Female	795-804	Female
795-804	Male	805-814	Male
805-814	Female	815-824	Female
815-824	Male	825-834	Male
825-834	Female	835-844	Female
835-844	Male	845-854	Male
845-854	Female	855-864	Female
855-864	Male	865-874	Male
865-874	Female	875-884	Female

<400> 37

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Variable	Mean	SD	Min	Max
Age	34.5	10.2	21	55
Gender				
Male	15.2	3.5	0	25
Female	19.8	4.1	0	30
Marital Status				
Married	12.5	2.8	0	20
Single	22.3	5.2	0	35
Divorced	1.2	0.5	0	3
Widowed	0.8	0.3	0	2
Education				
High School	8.5	2.1	0	12
College	15.2	3.5	0	25
Postgraduate	5.3	1.8	0	10
Income				
Low	10.5	2.5	0	20
Medium	18.2	4.2	0	30
High	7.3	1.9	0	15
Health Status				
Good	18.5	3.8	0	25
Fair	12.3	2.9	0	20
Poor	8.7	2.2	0	15
Stress Level				
Low	15.8	3.2	0	25
Medium	12.1	2.7	0	20
High	9.4	2.3	0	15
Life Satisfaction				
High	14.2	3.1	0	25
Medium	11.5	2.6	0	20
Low	8.9	2.1	0	15

<210> 42

<211> 243
 <212> PRT
 <213> Homo Sapien

<400> 42

Met	Arg	Pro	Leu	Leu	Val	Leu	Leu	Leu	Gly	Leu	Ala	Ala	Gly	
1				5					10				15	
Ser	Pro	Pro	Leu	Asp	Asp	Asn	Lys	Ile	Pro	Ser	Leu	Cys	Pro	Gly
				20					25					30
His	Pro	Gly	Leu	Pro	Gly	Thr	Pro	Gly	His	His	Gly	Ser	Gln	Gly
				35					40					45
Leu	Pro	Gly	Arg	Asp	Gly	Arg	Asp	Gly	Arg	Asp	Gly	Ala	Pro	Gly
				50					55					60
Ala	Pro	Gly	Glu	Lys	Gly	Glu	Gly	Gly	Arg	Pro	Gly	Leu	Pro	Gly
				65					70					75
Pro	Arg	Gly	Asp	Pro	Gly	Pro	Arg	Gly	Glu	Ala	Gly	Pro	Ala	Gly
				80					85					90
Pro	Thr	Gly	Pro	Ala	Gly	Glu	Cys	Ser	Val	Pro	Pro	Arg	Ser	Ala
				95					100					105
Phe	Ser	Ala	Lys	Arg	Ser	Glu	Ser	Arg	Val	Pro	Pro	Pro	Ser	Asp
				110					115					120
Ala	Pro	Leu	Pro	Phe	Asp	Arg	Val	Leu	Val	Asn	Glu	Gln	Gly	His
				125					130					135
Tyr	Asp	Ala	Val	Thr	Gly	Lys	Phe	Thr	Cys	Gln	Val	Pro	Gly	Val
				140					145					150
Tyr	Tyr	Phe	Ala	Val	His	Ala	Thr	Val	Tyr	Arg	Ala	Ser	Leu	Gln
				155					160					165
Phe	Asp	Leu	Val	Lys	Asn	Gly	Glu	Ser	Ile	Ala	Ser	Phe	Phe	Gln
				170					175					180
Phe	Phe	Gly	Gly	Trp	Pro	Lys	Pro	Ala	Ser	Leu	Ser	Gly	Gly	Ala
				185					190					195
Met	Val	Arg	Leu	Glu	Pro	Glu	Asp	Gln	Val	Trp	Val	Gln	Val	Gly
				200					205					210
Val	Gly	Asp	Tyr	Ile	Gly	Ile	Tyr	Ala	Ser	Ile	Lys	Thr	Asp	Ser
				215					220					225
Thr	Phe	Ser	Gly	Phe	Leu	Val	Tyr	Ser	Asp	Trp	His	Ser	Ser	Pro
				230					235					240
Val	Phe	Ala												

<210> 43
 <211> 24

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide probe

 <400> 43
 tacaggccca gtcaggacca gggg 24

 <210> 44
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide probe

 <400> 44
 agccagcctc gctctcgg 18

 <210> 45
 <211> 18
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide probe

 <400> 45
 gtctgcgatc aggtctgg 18

 <210> 46
 <211> 20
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide probe

 <400> 46
 gaaagaggca atggattcgc 20

 <210> 47
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> Synthetic oligonucleotide probe

 <400> 47
 gacttacact tgccagcaca gcac 24

 <210> 48
 <211> 45
 <212> DNA
 <213> Artificial Sequence

Variable	Mean	SD	Min	Max
Age (years)	34.5	10.2	18	65
Gender (Male/Female)	15/15			
Education (years)	12.5	1.5	8	16
Occupation (Student/Worker)	10/5			
Marital status (Single/Married)	12/3			
Family size (number of children)	1.5	1.0	0	4
Income (USD/month)	150	50	50	300
Health status (Healthy/Unhealthy)	12/3			
Smoking status (Smoker/Non-smoker)	5/10			
Alcohol consumption (Yes/No)	3/12			
Exercise frequency (times/week)	2.5	1.5	0	5
Dietary habits (Vegetarian/Non-vegetarian)	8/7			
Stress level (Low/Medium/High)	4/6/5			
Sleep quality (Good/Poor)	10/5			
Work-life balance (Satisfied/Unsatisfied)	7/8			
Overall life satisfaction (1-5)	3.5	0.8	1	5

<223> Synthetic oligonucleotide probe

<400> 48

ggagcaccac caactggagg gtccggagta gcgagcgccc cgaag 45

<210> 49

<211> 1876

<212> DNA

<213> Homo Sapien

<400> 49

ctcttttgtc caccagccca gcctgactcc tggagattgt gaatagctcc 50

atccagcctg agaaacaagc cgggtggctg agccaggctg tgcacggagc 100

acctgacggg cccaacagac ccattgctgca tccagagacc tcccctggcc 150

gggggcatct cctggctgtg ctctggccc tccttggcac cacctgggca 200

gaggtgtggc caccacagct gcaggagcag gctccgatgg ccggagccct 250

gaacaggaag gagaqtttct tgetcctctc cctgcacaac cgcctgcgca 300

gctgggtcca gccccctgcg gctgacatgc ggaggctgga ctggaagtgc 350

agcctggccc aactggctca agccaagggca gccctctgtg gaatcccaac 400

cccgagcctg gcatccggcc tgtggcgcac cctgcaagtg ggctggaaca 450

tgcagctgct gcccgcgggc ttggcgtcct ttgttgaagt ggtcagccta 500

tggtttgcag agggggcagcg gtacagccac gcggcaggaq agtqtgctcg 550

caacgccacc tgcacccact acacgcagct cgtgtgggcc acctcaagcc 600

agctgggctg tgggcggcac ctgtgctctg caggccagac agcgatagaa 650

gcctttgtct gtgcctactc ccccgqaggg aactgggagg tcaacqggaa 700

gacaatcatc ccctataaga aggggtgcctg gtgttcgctc tgcacagcca 750

gtgtctcagg ctgcttcaaa gcctgggacc atgcaggggg gctctgtgag 800

gtccccagga atccttgtcg catgaqctgc cagaaccatg qacgtctcaa 850

catcagcacc tgccactgcc actgtccccc tggctacacg ggcagatact 900

gccaaagtgag gtgcagcctg cagtgtgtgc acggccgggtt ccggggaggag 950

gagtgtctcgt gcgctctgtga catcggctac ggggggaagccc agtgtgtccac 1000

caaggtgcat tttcccttcc acacctgtga cctgaggatc gacggagact 1050

gcttcatggt gtcttcagag gcagacacct attacagagc caggatgaaa 1100

tgtcagagga aagqcqqqqt qctggcccaq atcaagagcc agaaaqtqca 1150

ggacatcctc gccttctatc tgggcccgcct ggagaccacc aacgaggtga 1200
 ctgacagtga cttcgagacc aggaacttct ggatcgggct cacctacaag 1250
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 cagttttgcc tttgggcagc ctgacaacca cgggctgggtg tggctgagtg 1350
 ctgccatggg gtttggcaac tgcgtggagc tgcaggcttc agctgccttc 1400
 aactggaacg accagcgctg caaaacccga aaccgttaca tctgccagtt 1450
 tgcccaggag cacatctccc ggtggggccc agggctcctga ggcttgacca 1500
 catggctccc tcgctgccc tgggagcacc ggctctgctt acctgtctgc 1550
 ccacctgtct ggaacaaggg ccagggttaag accacatgcc tcatgtccaa 1600
 agaggtctca gaccttgac aatgccagaa gttgggcaga gagaggcagg 1650
 gaggccagtg agggccaggg agtgagtgtt agaagaagct ggggcccttc 1700
 gcctgctttt gattgggaag atgggcttca attagatggc gaaggagagg 1750
 acaccgccag tgggtcaaaa aggctgctct cttccacctg gccagaccc 1800
 tgtggggcag cggagcttcc ctgtggcatg aacccacagg ggtattaaat 1850
 tatgaatcag ctgaaaaaaaa aaaaaa 1876

<210> 50

<211> 455

<212> PRT

<213> Homo Sapien

<400> 50

Met	Leu	His	Pro	Glu	Thr	Ser	Pro	Gly	Arg	Gly	His	Leu	Leu	Ala
1				5				10						15

Val	Leu	Leu	Ala	Leu	Leu	Gly	Thr	Thr	Trp	Ala	Glu	Val	Trp	Pro
				20					25					30

Pro	Gln	Leu	Gln	Glu	Gln	Ala	Pro	Met	Ala	Gly	Ala	Leu	Asn	Arg
				35					40					45

Lys	Glu	Ser	Phe	Leu	Leu	Leu	Ser	Leu	His	Asn	Arg	Leu	Arg	Ser
				50					55					60

Trp	Val	Gln	Pro	Pro	Ala	Ala	Asp	Met	Arg	Arg	Leu	Asp	Trp	Ser
				65					70					75

Asp	Ser	Leu	Ala	Gln	Leu	Ala	Gln	Ala	Arg	Ala	Ala	Leu	Cys	Gly
				80					85					90

Ile	Pro	Thr	Pro	Ser	Leu	Ala	Ser	Gly	Leu	Trp	Arg	Thr	Leu	Gln
				95					100					105

Val	Gly	Trp	Asn	Met	Gln	Leu	Leu	Pro	Ala	Gly	Leu	Ala	Ser	Phe
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

				110					115					120
Val	Glu	Val	Val	Ser 125	Leu	Trp	Phe	Ala	Glu 130	Gly	Gln	Arg	Tyr	Ser 135
His	Ala	Ala	Gly	Glu 140	Cys	Ala	Arg	Asn	Ala 145	Thr	Cys	Thr	His	Tyr 150
Thr	Gln	Leu	Val	Trp 155	Ala	Thr	Ser	Ser	Gln 160	Leu	Gly	Cys	Gly	Arg 165
His	Leu	Cys	Ser	Ala 170	Gly	Gln	Thr	Ala	Ile 175	Glu	Ala	Phe	Val	Cys 180
Ala	Tyr	Ser	Pro	Gly 185	Gly	Asn	Trp	Glu	Val 190	Asn	Gly	Lys	Thr	Ile 195
Ile	Pro	Tyr	Lys	Lys 200	Gly	Ala	Trp	Cys	Ser 205	Leu	Cys	Thr	Ala	Ser 210
Val	Ser	Gly	Cys	Phe 215	Lys	Ala	Trp	Asp	His 220	Ala	Gly	Gly	Leu	Cys 225
Glu	Val	Pro	Arg	Asn 230	Pro	Cys	Arg	Met	Ser 235	Cys	Gln	Asn	His	Gly 240
Arg	Leu	Asn	Ile	Ser 245	Thr	Cys	His	Cys	His 250	Cys	Pro	Pro	Gly	Tyr 255
Thr	Gly	Arg	Tyr	Cys 260	Gln	Val	Arg	Cys	Ser 265	Leu	Gln	Cys	Val	His 270
Gly	Arg	Phe	Arg	Glu 275	Glu	Glu	Cys	Ser	Cys 280	Val	Cys	Asp	Ile	Gly 285
Tyr	Gly	Gly	Ala	Gln 290	Cys	Ala	Thr	Lys	Val 295	His	Phe	Pro	Phe	His 300
Thr	Cys	Asp	Leu	Arg 305	Ile	Asp	Gly	Asp	Cys 310	Phe	Met	Val	Ser	Ser 315
Glu	Ala	Asp	Thr	Tyr 320	Tyr	Arg	Ala	Arg	Met 325	Lys	Cys	Gln	Arg	Lys 330
Gly	Gly	Val	Leu	Ala 335	Gln	Ile	Lys	Ser	Gln 340	Lys	Val	Gln	Asp	Ile 345
Leu	Ala	Phe	Tyr	Leu 350	Gly	Arg	Leu	Glu	Thr 355	Thr	Asn	Glu	Val	Thr 360
Asp	Ser	Asp	Phe	Glu 365	Thr	Arg	Asn	Phe	Trp 370	Ile	Gly	Leu	Thr	Tyr 375
Lys	Thr	Ala	Lys	Asp 380	Ser	Phe	Arg	Trp	Ala 385	Thr	Gly	Glu	His	Gln 390
Ala	Phe	Thr	Ser	Phe 395	Ala	Phe	Gly	Gln	Pro 400	Asp	Asn	His	Gly	Leu 405

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<211> 694

<212> PRT

<213> Homo Sapien

<400> 55

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Gly	Ser	Gln	Glu	Glu	Ala	Gln	Ser	Trp	Gly	His	Ser	Ser	Glu	Gln
			20						25					30
Asp	Gly	Leu	Arg	Val	Pro	Arg	Gln	Val	Arg	Leu	Leu	Gln	Arg	Leu
			35						40					45
Lys	Thr	Lys	Pro	Leu	Met	Thr	Glu	Phe	Ser	Val	Lys	Ser	Thr	Ile
			50						55					60
Ile	Ser	Arg	Tyr	Ala	Phe	Thr	Thr	Val	Ser	Cys	Arg	Met	Leu	Asn
			65						70					75
Arg	Ala	Ser	Glu	Asp	Gln	Asp	Ile	Glu	Phe	Gln	Met	Gln	Ile	Pro
			80						85					90
Ala	Ala	Ala	Phe	Ile	Thr	Asn	Phe	Thr	Met	Leu	Ile	Gly	Asp	Lys
			95						100					105
Val	Tyr	Gln	Gly	Glu	Ile	Thr	Glu	Arg	Glu	Lys	Lys	Ser	Gly	Asp
			110						115					120

Arg Val Lys Glu Lys Arg Asn Lys Thr Thr Glu Glu Asn Gly Glu	125	130	135
Lys Gly Thr Glu Ile Phe Arg Ala Ser Ala Val Ile Pro Ser Lys	140	145	150
Asp Lys Ala Ala Phe Phe Leu Ser Tyr Glu Glu Leu Leu Gln Arg	155	160	165
Arg Leu Gly Lys Tyr Glu His Ser Ile Ser Val Arg Pro Gln Gln	170	175	180
Leu Ser Gly Arg Leu Ser Val Asp Val Asn Ile Leu Glu Ser Ala	185	190	195
Gly Ile Ala Ser Leu Glu Val Leu Pro Leu His Asn Ser Arg Gln	200	205	210
Arg Gly Ser Gly Arg Gly Glu Asp Asp Ser Gly Pro Pro Pro Ser	215	220	225
Thr Val Ile Asn Gln Asn Glu Thr Phe Ala Asn Ile Ile Phe Lys	230	235	240
Pro Thr Val Val Gln Gln Ala Arg Ile Ala Gln Asn Gly Ile Leu	245	250	255
Gly Asp Phe Ile Ile Arg Tyr Asp Val Asn Arg Glu Gln Ser Ile	260	265	270
Gly Asp Ile Gln Val Leu Asn Gly Tyr Phe Val His Tyr Phe Ala	275	280	285
Pro Lys Asp Leu Pro Pro Leu Pro Lys Asn Val Val Phe Val Leu	290	295	300
Asp Ser Ser Ala Ser Met Val Gly Thr Lys Leu Arg Gln Thr Lys	305	310	315
Asp Ala Leu Phe Thr Ile Leu His Asp Leu Arg Pro Gln Asp Arg	320	325	330
Phe Ser Ile Ile Gly Phe Ser Asn Arg Ile Lys Val Trp Lys Asp	335	340	345
His Leu Ile Ser Val Thr Pro Asp Ser Ile Arg Asp Gly Lys Val	350	355	360
Tyr Ile His His Met Ser Pro Thr Gly Gly Thr Asp Ile Asn Gly	365	370	375
Ala Leu Gln Arg Ala Ile Arg Leu Leu Asn Lys Tyr Val Ala His	380	385	390
Ser Gly Ile Gly Asp Arg Ser Val Ser Leu Ile Val Phe Leu Thr	395	400	405
Asp Gly Lys Pro Thr Val Gly Glu Thr His Thr Leu Lys Ile Leu			

410	415	420
Asn Asn Thr Arg Glu Ala Ala Arg Gly Gln Val Cys Ile Phe Thr		
425	430	435
Ile Gly Ile Gly Asn Asp Val Asp Phe Arg Leu Leu Glu Lys Leu		
440	445	450
Ser Leu Glu Asn Cys Gly Leu Thr Arg Arg Val His Glu Glu Glu		
455	460	465
Asp Ala Gly Ser Gln Leu Ile Gly Phe Tyr Asp Glu Ile Arg Thr		
470	475	480
Pro Leu Leu Ser Asp Ile Arg Ile Asp Tyr Pro Pro Ser Ser Val		
485	490	495
Val Gln Ala Thr Lys Thr Leu Phe Pro Asn Tyr Phe Asn Gly Ser		
500	505	510
Glu Ile Ile Ile Ala Gly Lys Leu Val Asp Arg Lys Leu Asp His		
515	520	525
Leu His Val Glu Val Thr Ala Ser Asn Ser Lys Lys Phe Ile Ile		
530	535	540
Leu Lys Thr Asp Val Pro Val Arg Pro Gln Lys Ala Gly Lys Asp		
545	550	555
Val Thr Gly Ser Pro Arg Pro Gly Gly Asp Gly Glu Gly Asp Thr		
560	565	570
Asn His Ile Glu Arg Leu Trp Ser Tyr Leu Thr Thr Lys Glu Leu		
575	580	585
Leu Ser Ser Trp Leu Gln Ser Asp Asp Glu Pro Glu Lys Glu Arg		
590	595	600
Leu Arg Gln Arg Ala Gln Ala Leu Ala Val Ser Tyr Arg Phe Leu		
605	610	615
Thr Pro Phe Thr Ser Met Lys Leu Arg Gly Pro Val Pro Arg Met		
620	625	630
Asp Gly Leu Glu Glu Ala His Gly Met Ser Ala Ala Met Gly Pro		
635	640	645
Glu Pro Val Val Gln Ser Val Arg Gly Ala Gly Thr Gln Pro Gly		
650	655	660
Pro Leu Leu Lys Lys Pro Asn Ser Val Lys Lys Lys Gln Asn Lys		
665	670	675
Thr Lys Lys Arg His Gly Arg Asp Gly Val Phe Pro Leu His His		
680	685	690
Leu Gly Ile Arg		

<210> 56
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 56
gtgggaacca aactccggca gacc 24

<210> 57
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 57
cacatcgagc gtctctgg 18

<210> 58
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 58
agccgctcct tctccggttc atcg 24

<210> 59
<211> 48
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 59
tggaaggacc acttgatc agtcactcca gacagcatca gggatggg 48

<210> 60
<211> 1413
<212> DNA
<213> Homo Sapien

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tgctgttgct cttctccgcc gcggcactga tccccacagg tgatgggcag 150
aatctgttta cgaaagacgt gacagtgatc gagggagagg ttgcgaccat 200



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acagtgatat tgg 1413

<210> 61

<211> 440

<212> PRT

<213> Homo Sapien

<400> 61

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320	325	330
Pro Pro Thr Thr Ile Pro Pro Pro Thr Thr Thr Thr Thr Thr		
335	340	345
Thr Thr Thr Thr Thr Thr Ile Leu Thr Ile Ile Thr Asp Ser Arg		
350	355	360
Ala Gly Glu Glu Gly Ser Ile Arg Ala Val Asp His Ala Val Ile		
365	370	375
Gly Gly Val Val Ala Val Val Val Phe Ala Met Leu Cys Leu Leu		
380	385	390
Ile Ile Leu Gly Arg Tyr Phe Ala Arg His Lys Gly Thr Tyr Phe		
395	400	405
Thr His Glu Ala Lys Gly Ala Asp Asp Ala Ala Asp Ala Asp Thr		
410	415	420
Ala Ile Ile Asn Ala Glu Gly Gly Gln Asn Asn Ser Glu Glu Lys		
425	430	435
Lys Glu Tyr Phe Ile		
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<210> 62
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 62
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<210> 63
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 63
 gtacactgtg accagtcagc 20

<210> 64
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

<400> 64
atcatcacag attcccgagc 20

<210> 65
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 65
ttcaatctcc tcaccttcca ccgc 24

<210> 66
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<220>
<223> Synthetic oligonucleotide probe

<400> 66
atagctgtgt ctgcgtctgc tgcg 24

<210> 67
<211> 50
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 67
cgcggcactg atccccacag gtgatgggca gaatctgttt acgaaagacg 50

<210> 68
<211> 2555
<212> DNA
<213> Homo Sapien

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<210> 69
<211> 598
<212> PRT
<213> Homo Sapien

<400> 69
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20 25 30
Ser Gln Pro Gln Thr Val Phe Cys Thr Ala Arg Gln Gly Thr Thr
35 40 45
Val Pro Arg Asp Val Pro Pro Asp Thr Val Gly Leu Tyr Val Phe
50 55 60
Glu Asn Gly Ile Thr Met Leu Asp Ala Ser Ser Phe Ala Gly Leu
65 70 75
Pro Gly Leu Gln Leu Leu Asp Leu Ser Gln Asn Gln Ile Ala Ser
80 85 90
Leu Arg Leu Pro Arg Leu Leu Leu Leu Asp Leu Ser His Asn Ser
95 100 105
Leu Leu Ala Leu Glu Pro Gly Ile Leu Asp Thr Ala Asn Val Glu

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Ala	Leu	Arg	Leu	Ala 125	Gly	Leu	Gly	Leu	Gln 130	Gln	Leu	Asp	Glu	Gly 135
Leu	Phe	Ser	Arg	Leu 140	Arg	Asn	Leu	His	Asp 145	Leu	Asp	Val	Ser	Asp 150
Asn	Gln	Leu	Glu	Arg 155	Val	Pro	Pro	Val	Ile 160	Arg	Gly	Leu	Arg	Gly 165
Leu	Thr	Arg	Leu	Arg 170	Leu	Ala	Gly	Asn	Thr 175	Arg	Ile	Ala	Gln	Leu 180
Arg	Pro	Glu	Asp	Leu 185	Ala	Gly	Leu	Ala	Ala 190	Leu	Gln	Glu	Leu	Asp 195
Val	Ser	Asn	Leu	Ser 200	Leu	Gln	Ala	Leu	Pro 205	Gly	Asp	Leu	Ser	Gly 210
Leu	Phe	Pro	Arg	Leu 215	Arg	Leu	Leu	Ala	Ala 220	Ala	Arg	Asn	Pro	Phe 225
Asn	Cys	Val	Cys	Pro 230	Leu	Ser	Trp	Phe	Gly 235	Pro	Trp	Val	Arg	Glu 240
Ser	His	Val	Thr	Leu 245	Ala	Ser	Pro	Glu	Glu 250	Thr	Arg	Cys	His	Phe 255
Pro	Pro	Lys	Asn	Ala 260	Gly	Arg	Leu	Leu	Leu 265	Glu	Leu	Asp	Tyr	Ala 270
Asp	Phe	Gly	Cys	Pro 275	Ala	Thr	Thr	Thr	Thr 280	Ala	Thr	Val	Pro	Thr 285
Thr	Arg	Pro	Val	Val 290	Arg	Glu	Pro	Thr	Ala 295	Leu	Ser	Ser	Ser	Leu 300
Ala	Pro	Thr	Trp	Leu 305	Ser	Pro	Thr	Ala	Pro 310	Ala	Thr	Glu	Ala	Pro 315
Ser	Pro	Pro	Ser	Thr 320	Ala	Pro	Pro	Thr	Val 325	Gly	Pro	Val	Pro	Gln 330
Pro	Gln	Asp	Cys	Pro 335	Pro	Ser	Thr	Cys	Leu 340	Asn	Gly	Gly	Thr	Cys 345
His	Leu	Gly	Thr	Arg 350	His	His	Leu	Ala	Cys 355	Leu	Cys	Pro	Glu	Gly 360
Phe	Thr	Gly	Leu	Tyr 365	Cys	Glu	Ser	Gln	Met 370	Gly	Gln	Gly	Thr	Arg 375
Pro	Ser	Pro	Thr	Pro 380	Val	Thr	Pro	Arg	Pro 385	Pro	Arg	Ser	Leu	Thr 390
Leu	Gly	Ile	Glu	Pro 395	Val	Ser	Pro	Thr	Ser 400	Leu	Arg	Val	Gly	Leu 405

<400> 71
cggttctgagg gacgttaggg ctcg 24

<210> 72
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 72
ctgcccaccg tccacctgcc tcaat 25

<210> 73
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 73
aggactgccc accgtccacc tgcctcaatg ggggcacatg ccacc 45

<210> 74
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic Oligonucleotide Probe

<400> 74
acgcaaagcc ctacatctaa gccagagaga gacagggcag ctggg 45

<210> 75
<211> 1077
<212> DNA
<213> Homo Sapien

<400> 75
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ggcctccagg caacatgggg ggcccagtca gagagccggc actctcagtt 200
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gccggctgca ggggacagga ggccctccc agaatgggga agggatatccc 350
tggcagagtc tcccggagca gagttccgat gccctggaag cctggggagaa 400

Val	Pro	Ile	Asn	Ala	Thr	Ser	Lys	Asp	Asp	Ser	Asp	Val	Thr	Glu
			125						130					135
Val	Met	Trp	Gln	Pro	Ala	Leu	Arg	Arg	Gly	Arg	Gly	Leu	Gln	Ala
			140						145					150
Gln	Gly	Tyr	Gly	Val	Arg	Ile	Gln	Asp	Ala	Gly	Val	Tyr	Leu	Leu
			155						160					165
Tyr	Ser	Gln	Val	Leu	Phe	Gln	Asp	Val	Thr	Phe	Thr	Met	Gly	Gln
			170						175					180
Val	Val	Ser	Arg	Glu	Gly	Gln	Gly	Arg	Gln	Glu	Thr	Leu	Phe	Arg
			185						190					195
Cys	Ile	Arg	Ser	Met	Pro	Ser	His	Pro	Asp	Arg	Ala	Tyr	Asn	Ser
			200						205					210
Cys	Tyr	Ser	Ala	Gly	Val	Phe	His	Leu	His	Gln	Gly	Asp	Ile	Leu
			215						220					225
Ser	Val	Ile	Ile	Pro	Arg	Ala	Arg	Ala	Lys	Leu	Asn	Leu	Ser	Pro
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His	Gly	Thr	Phe	Leu	Gly	Phe	Val	Lys	Leu					
			245					250						

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 <212> DNA
 <213> Homo Sapien

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<210> 78
 <211> 281
 <212> PRT
 <213> Homo Sapien

<400> 78
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 35 40 45
 Pro Pro Asp His Ala Glu Arg Ala Glu Glu Gln His Glu Lys Tyr
 50 55 60
 Arg Pro Ser Gln Asp Gln Gly Leu Pro Ala Ser Arg Cys Leu Arg
 65 70 75
 Cys Cys Asp Pro Gly Thr Ser Met Tyr Pro Ala Thr Ala Val Pro
 80 85 90
 Gln Ile Asn Ile Thr Ile Leu Lys Gly Glu Lys Gly Asp Arg Gly
 95 100 105
 Asp Arg Gly Leu Gln Gly Lys Tyr Gly Lys Thr Gly Ser Ala Gly

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125	130	135
Ala Pro Gly Glu Arg Cys Lys Ser His	Tyr Ala Ala Phe Ser Val	
140	145	150
Gly Arg Lys Lys Pro Met His Ser Asn	His Tyr Tyr Gln Thr Val	
155	160	165
Ile Phe Asp Thr Glu Phe Val Asn Leu	Tyr Asp His Phe Asn Met	
170	175	180
Phe Thr Gly Lys Phe Tyr Cys Tyr Val	Pro Gly Leu Tyr Phe Phe	
185	190	195
Ser Leu Asn Val His Thr Trp Asn Gln	Lys Glu Thr Tyr Leu His	
200	205	210
Ile Met Lys Asn Glu Glu Glu Val Val	Ile Leu Phe Ala Gln Val	
215	220	225
Gly Asp Arg Ser Ile Met Gln Ser Gln	Ser Leu Met Leu Glu Leu	
230	235	240
Arg Glu Gln Asp Gln Val Trp Val Arg	Leu Tyr Lys Gly Glu Arg	
245	250	255
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260	265	270
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 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic oligonucleotide probe

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<210> 80
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 <212> DNA
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<220>
 <223> Synthetic oligonucleotide probe

<400> 80
 ctgaagaagt agaggccggg cacg 24

<210> 81

<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 81
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<210> 82
<211> 2284
<212> DNA
<213> Homo Sapien

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cttcttaaag caaactaaga ccagagggag gattatcctt gacctttgaa 200
gacaaaaact aaactgaaat ttaaaatgtt cttcggggga gaaggagct 250
tgacttacac tttggtaata atttgcttcc tgacactaag gctgtctgct 300
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Ser	Lys	Gly	Ile	Arg 50	Gly	Asn	Glu	Pro	Val 55	Tyr	Thr	Ser	Thr	Gln 60
Glu	Asp	Cys	Ile	Asn 65	Ser	Cys	Cys	Ser	Thr 70	Lys	Asn	Ile	Ser	Gly 75
Asp	Lys	Ala	Cys	Asn 80	Leu	Met	Ile	Phe	Asp 85	Thr	Arg	Lys	Thr	Ala 90
Arg	Gln	Pro	Asn	Cys 95	Tyr	Leu	Phe	Phe	Cys 100	Pro	Asn	Glu	Glu	Ala 105
Cys	Pro	Leu	Lys	Pro 110	Ala	Lys	Gly	Leu	Met 115	Ser	Tyr	Arg	Ile	Ile 120
Thr	Asp	Phe	Pro	Ser 125	Leu	Thr	Arg	Asn	Leu 130	Pro	Ser	Gln	Glu	Leu 135
Pro	Gln	Glu	Asp	Ser 140	Leu	Leu	His	Gly	Gln 145	Phe	Ser	Gln	Ala	Val 150
Thr	Pro	Leu	Ala	His 155	His	His	Thr	Asp	Tyr 160	Ser	Lys	Pro	Thr	Asp 165
Ile	Ser	Trp	Arg	Asp 170	Thr	Leu	Ser	Gln	Lys 175	Phe	Gly	Ser	Ser	Asp 180
His	Leu	Glu	Lys	Leu 185	Phe	Lys	Met	Asp	Glu 190	Ala	Ser	Ala	Gln	Leu 195
Leu	Ala	Tyr	Lys	Glu 200	Lys	Gly	His	Ser	Gln 205	Ser	Ser	Gln	Phe	Ser 210
Ser	Asp	Gln	Glu	Ile 215	Ala	His	Leu	Leu	Pro 220	Glu	Asn	Val	Ser	Ala 225
Leu	Pro	Ala	Thr	Val 230	Ala	Val	Ala	Ser	Pro 235	His	Thr	Thr	Ser	Ala 240
Thr	Pro	Lys	Pro	Ala 245	Thr	Leu	Leu	Pro	Thr 250	Asn	Ala	Ser	Val	Thr 255
Pro	Ser	Gly	Thr	Ser 260	Gln	Pro	Gln	Leu	Ala 265	Thr	Thr	Ala	Pro	Pro 270
Val	Thr	Thr	Val	Thr 275	Ser	Gln	Pro	Pro	Thr 280	Thr	Leu	Ile	Ser	Thr 285
Val	Phe	Thr	Arg	Ala 290	Ala	Ala	Thr	Leu	Gln 295	Ala	Met	Ala	Thr	Thr 300
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<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 87
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<210> 88
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 88
agtgtagtc aagctccc 18

<210> 89
<211> 49
<212> DNA
<213> Artificial Sequence

<220>
<223> Synthetic oligonucleotide probe

<400> 89
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<210> 90
<211> 957
<212> DNA
<213> Homo Sapien

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cattccagat gcaccctgt ccagtgtgc ctatagcatc cgcagcatcg 150
gggagaggcc tgtcctcaaa gctccagtcc ccaaaaggca aaaatgtgac 200
cactggactc cctgcccatac tgacacctat gcctacaggt tactcagcgg 250
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tgggagaaca gctgggaaat gttgccagag gaataaacat tgccattgtc 350
aactatgtaa ctgggaatgt gacagcaaca cgatgttttg atatgtatga 400
aggcgataac tctggaccga tgacaaagtt tattcagagt gctgctccaa 450
aatccctgct cttcatgggtg acctatgacg acggaagcac aagactgaat 500

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<400> 91

65

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Asn	Met	Lys	Phe	Arg	Ser	Ser	Trp	Val	Phe	Ile	Ala	Ala	Lys	Gly
			185						190					195
Leu	Glu	Leu	Pro	Ser	Glu	Ile	Gln	Arg	Glu	Lys	Ile	Asn	His	Ser
			200						205					210
Asp	Ala	Lys	Asn	Asn	Arg	Tyr	Ser	Gly	Trp	Pro	Ala	Glu	Ile	Gln
			215						220					225
Ile	Glu	Gly	Cys	Ile	Pro	Lys	Glu	Arg	Ser					
			230						235					

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<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

<400> 92

aatgtgacca ctggactccc 20

<210> 93

<211> 18

<212> DNA

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<220>

<223> Synthetic oligonucleotide probe

<400> 93

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<210> 94

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

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<210> 95

<211> 47

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

<400> 95

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<210> 96
 <211> 21
 <212> DNA
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 <220>
 <223> Synthetic oligonucleotide probe

 <400> 96
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 <210> 97
 <211> 25
 <212> DNA
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 <220>
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 <400> 97
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 <210> 98
 <211> 18
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 <220>
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 <400> 98
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